## Fuduan Peng

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9129130/publications.pdf

Version: 2024-02-01

1040056 1199594 12 390 9 12 citations h-index g-index papers 13 13 13 799 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Populationâ€scale genetic control of alternative polyadenylation and its association with human diseases. Quantitative Biology, 2022, 10, 44-54.	0.5	2
2	A genomeâ€wide association study identifies <scp><i>FSHR</i></scp> rs2300441 associated with follicleâ€stimulating hormone levels. Clinical Genetics, 2020, 97, 869-877.	2.0	8
3	Validation of methylation-based forensic age estimation in time-series bloodstains on FTA cards and gauze at room temperature conditions. Forensic Science International: Genetics, 2019, 40, 168-174.	3.1	11
4	Genome-Wide Association Studies Identify MultipleÂGenetic Loci Influencing Eyebrow ColorÂVariation in Europeans. Journal of Investigative Dermatology, 2019, 139, 1601-1605.	0.7	17
5	A landscape of synthetic viable interactions in cancer. Briefings in Bioinformatics, 2018, 19, bbw142.	<b>6.</b> 5	9
6	Meta-analysis of genome-wide association studies identifies 8 novel loci involved in shape variation of human head hair. Human Molecular Genetics, 2018, 27, 559-575.	2.9	51
7	Systematic feature selection improves accuracy of methylation-based forensic age estimation in Han Chinese males. Forensic Science International: Genetics, 2018, 35, 38-45.	3.1	46
8	Genome-wide association studies and CRISPR/Cas9-mediated gene editing identify regulatory variants influencing eyebrow thickness in humans. PLoS Genetics, 2018, 14, e1007640.	<b>3.</b> 5	20
9	EMT-Regulome: a database for EMT-related regulatory interactions, motifs and network. Cell Death and Disease, 2017, 8, e2872-e2872.	6.3	21
10	Differential expression analysis at the individual level reveals a lncRNA prognostic signature for lung adenocarcinoma. Molecular Cancer, 2017, 16, 98.	19.2	101
11	Autophagyâ€related prognostic signature for breast cancer. Molecular Carcinogenesis, 2016, 55, 292-299.	2.7	68
12	The BRCA1/2-directed miRNA signature predicts a good prognosis in ovarian cancer patients with wild-type BRCA1/2. Oncotarget, 2015, 6, 2397-2406.	1.8	36